

Fig. 1

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A. t. 6 SAPGKVLMTGGYLVLEKPNAGLVLTSTNARFYAIVKPINEEVKPESWAWKW 55
      ||||| |: |||||: |. .|| |:| | :.
S. c. 8 SAPGKALLAGGYLVLDTKYEAFVVGLSARMHAHAHPYGSLOQSDKF.... 53
      . . . . .

56 TDVKLTSPQL.SRESMYKLSLNHLTLQSVSASDSRNPFEHAIQYAIAAA 104
   :|:. | | :| :| : || |:|||:| | | .
54 .EVRVKSKQFKDGEWLYHISPKSGFI.PVSIGGSKNPFIEKVI..ANVFS 99
      . . . . .

105 HLATEKDKESSLHKLQLGLDITILGSNDFYSYRNQIESAGLPLTPESLGT 154
   : | | . :|| |. | . | :| | |
100 YFKPNMDDYCNRNLFV..IDIF...SDD..AYHSQEDS.....VTEHRG. 136
      . . . . .

155 LAPFASITFNAAESNGANSKPEVAKTGLGSSAAMTTAVVAALLHYLGVD 204
      | | .. || ||||| ||| : | . || : | |
137 .....NRRLSFHSHRIEEVPKTGLGSSAGLVTVLTTALASFF.VSD 176
      . . . . .

205 LSDPCKEGKFGCSLDVIHMIAQTSHCLAQGKVGSGFDVSCAVYGSQRYV 254
   | . . : :||| :|| .|| |||||:||||||. | ||| ||
177 LENNVDKYR.....EVIHNLAQVAHCQAQKGIGSGFDVAAARYGSIRYR 220
      . . . . .

255 RFSPEVLSFAQVAVTGLPLNEVIGTILKGKWDNKRTEFSLPPLMNLFLGE 304
   || | .:| . . . :. | . || : |.:|:
221 RFPPALISNLPDIGSATYGSKLHLVDEEDWNITIKSNHLPGLTLWMGD 270
      . . . . .

305 PGSGGSSTPSMVGAVKKWQMSDPEKARENWQNLSANLELETKLNDLSKL 354
      ||| | :| || | | . . : | || |. | :|
271 I.KNGSETVQLVQKVKNWYDSHPESLKIYTELDHANSRFGDGLSKLDRL 319
      . . . . .

355 AKDHWDVYLRVIKSC..SVLTSEKWVLHATEPINEAIIKELLEAREAMLR 402
   . | | .: .| . | :| :| | :| |.
320 HETHDDYSDQIFESLERNDCTCQKY.....PEITEVRDAVAT 356
      . . . . .

403 IRILMRQMGEAASVPIEPESQTQLLDSTMSAEGVLLAGVPGAGGFDAIFA 452
   || |.. . ||| || ||| . .||| :|||||:|||
357 IRRSFRKITKESGADIEPPVQTSLLDDCQTLKGVLTCLIPGAGGYDAIAV 406
      . . . . .

453 ITLGD 457
      || |
407 ITKOD 411

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